SEQUENCE LISTING

<110> Beraud, Christophe

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Craven, Andrew
     Yu, Ming
     Sakowicz, Roman
     Patel, Umesh A.
     Davies, Katherine A.
<120> NOVEL MOTOR PROTEINS AND METHODS FOR THEIR USE
<130> 020552-001410US
<140> US 09/883096
<141> 2001-06-15
<150> US 09/594,655
<151> 2000-06-15
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<223> Nucleic acid sequence of human kinesin motor
     protein gene HsKip3a (Figure 1).
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<223> Description of Artificial Sequence: HsKip3a gene.
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caccacagtg atggcagtgg aggacagcac gctgcaagta gtggtacggg tgcggccccc 180
cacccetegg gagetggaca gteageggeg geeagtggtt caggtggtgg acgagegggt 240
gctggtgttt aaccetgagg agcccgatgg agggttccct ggcctgaaat ggggtggcac 300
ccatgatggc cccaagaaga agggcaaaga cctgacgttt gtctttgacc gggtctttgg 360
cgaggcggcc acccaacagg acgtgttcca gcacaccacg cacagcgtcc tggacagctt 420
cctccagggc tacaactgct cagtgtttgc ctacggggcc accggggctg ggaagacaca 480
caccatgetg ggaagggagg gggaccccgg catcatgtac ctgaccaccg tggaactgta 540
caggegeetg gaggeeegee ageaggagaa geaettegag gtgeteatea getaeeagga 600
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ggaccccgac aagggggtgg tggtgcaagg actttctttc caccagccag cctcagccga 720
gcagctgctg gagatactga ccagggggaa ccgtaaccgc acgcagcacc ccactgatgc 780
caacgcgact tecteceget cecatgccat ettecagate tttgtgaage ageaggaceg 840
ggttccagga ctgacccagg ctgtccaggt ggccaagatg agcctgattg acctggctgg 900
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caaccgctct ctgctggcgc tcatcaacgt cctcaatgcc ttggccgatg caaagggccg 1020
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20.00

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gcagcccaag ccagtcgtgg gccacttctc agcacgggaa ctggatgggg accgttctaa 1680
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caggogocca ccaccacgoc cggctaattt tttgtatttt tagtagagat ggggtttcac 3540
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cagtcctggt gaagatgtgg atgataatgg tgccttgatt tccaaatgaa gacagcttta 3900
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<210> 2
<211> 864
<212> PRT
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<213> Artificial Sequence

<220>

<223> Amino acid sequence encoded by human kinesin motor protein gene HsKip3a (Figure 1).

<220>

<223> Description of Artificial Sequence: Amino acid sequence of HsKip3a.

Met 1	Ala	Val	Glu	Asp 5	Ser	Thr	Leu	GIn	10	Val	Val	Arg	vaı	15	Pro
Pro	Thr	Pro	Arg 20	Glu	Leu	Asp	Ser	Gln 25	Arg	Arg	Pro	Val	Val 30	Gln	Val
Val	Asp	Glu 35	Arg	Val	Leu	Val	Phe 40	Asn	Pro	Glu	Glu	Pro 45	qaA	Gly	Gly
Phe	Pro 50	Gly	Leu	Lys.	Trp	Gly 55	Gly	Thr	His	·Asp	Gly 60	Pro	Lys	Lys	Lys
Gly 65	Lys	Asp	Leu	Thr	Phe 70	Val	Phe	Asp	Arg	Val 75	Phe	Gly	Glu	Ala	Alá 80
Thr	Gln	Gln	qaA	Val 85	Phe	Gln	His	Thr	Thr 90	His	Ser	Val	Leu	Asp 95	Ser
Phe	Leu	Gln	Gly 100	Tyr	Asn	Сув	Ser	Val 105	Phe	Ala	Tyr	Gly	Ala 110	Thr	Gly
Ala	Gly	Lys 115	Thr	His	Thr	Met	Leu 120	Gly	Arg	Glu	Gly	Asp 125	Pro	Gly	Ile
Met	Tyr 130	Leu	Thr	Thr	Val	Glu 135	Leu	Tyr	Arg	Arg	Leu 140	Glu	Ala	Arg	Gln
Gln 145	Glu	Lys	His	Phe	Glu 150	Val	Leu	Ile	Ser	Tyr 155	Gln	Glu	Val	Tyr	Asn 160
Glu	Gln	Ile	His	Asp 165	Leu	Leu	Glu	Pro	Lys 170	Gly	Pro	Leu	Ala	Ile 175	Arg
Glu	Asp	Pro	Asp 180	Lys	Gly	Val	Val	Val 185	Gln	Gly	Leu	Ser	Phe 190	His	Gln
Pro	Ala	Ser 195	Ala	Glu	Gln	Leu	Leu 200	Glu	Ile	Leu	Thr	Arg 205	Gly	Asn	Arg
	210			His		215					220				
His 225	Ala	Ile	Phe	Gln	Ile 230	Phe	Val	Lys	Gln	Gln 235	Asp	Arg	Val	Pro	Gly 240
Leu	Thr	Gln	Ala	Val 245	Gln	Val	Ala	Lys	Met 250	Ser	Leu	Ile	Asp	Leu 255	Ala
Gly	Ser	Glu	Arg 260	Ala	Ser	Ser	Thr	His 265	Ala	ГÀв	Gly	Glu	Arg 270	Leu	Arg
Glu	Gly	Ala 275	Asn	Ile	Asn	Arg	Ser 280	Leu	Leu	Ala	Leu	Ile 285	Asn	Val	Leu
Asn	Ala 290	Leu	Ala	qaA	Ala	Lys 295	Gly	Arg	Lys	Thr	His 300	Val	Pro	Tyr	Arg
Asp 305	Ser	Lys	Leu	Thr	Arg 310	Leu	Leu	ŗÀa	Asp	Ser 315	Leu	Gly	Gly	Asn	Cys 320

Arg Thr Val Met Ile Ala Ala Ile Ser Pro Ser Ser Leu Thr Tyr Glu 335

Asp Thr Tyr Asn Thr Leu Lys Tyr Ala Asp Arg Ala Lys Glu Ile Arg 340

Leu Ser Leu Lys Ser Asn Val Thr Ser Leu Asp Cys His Ile Ser Gln

Leu Ser Leu Lys Ser Asn Val Thr Ser Leu Asp Cys His Ile Ser Gln 355 360 365

Tyr Ala Thr Ile Cys Gln Gln Leu Gln Ala Glu Val Ala Ala Leu Arg 370 375 380

Lys Lys Leu Gln Val Tyr Glu Gly Gly Gly Gln Pro Pro Pro Gln Asp 385 390 395 400

Leu Pro Gly Ser Pro Lys Ser Gly Pro Pro Pro Glu His Leu Pro Ser 405 410 415

Ser Pro Leu Pro Pro His Pro Pro Ser Gln Pro Cys Thr Pro Glu Leu 420 425 430

Pro Ala Gly Pro Arg Ala Leu Gln Glu Glu Ser Leu Gly Met Glu Ala 435 440 445

Gln Val Glu Arg Ala Met Glu Gly Asn Ser Ser Asp Gln Glu Gln Ser 450 455 460

Pro Glu Asp Glu Asp Glu Gly Pro Ala Glu Glu Val Pro Thr Gln Met 465 470 475 480

Pro Glu Gln Asn Pro Thr His Ala Leu Pro Glu Ser Pro Arg Leu Thr
485 490 495

Leu Gln Pro Lys Pro Val Val Gly His Phe Ser Ala Arg Glu Leu Asp 500 505 510

Gly Asp Arg Ser Lys Gln Leu Ala Leu Lys Val Leu Cys Val Ala Gln 515 520 525

Arg Gln Tyr Ser Leu Leu Gln Ala Ala Asn Leu Leu Thr Pro Asp Met 530 535 540

Ile Thr Glu Phe Glu Thr Leu Gln Gln Leu Val Gln Glu Glu Lys Ile 545 550 555 560

Glu Pro Gly Ala Glu Ala Leu Arg Thr Ser Gly Leu Ala Arg Gly Ala
565 570 575

Pro Leu Ala Gln Glu Leu Cys Ser Glu Ser Ile Pro Val Pro Ser Pro 580 585 590

Leu Cys Pro Glu Pro Pro Gly Tyr Thr Gly Pro Val Thr Arg Thr Met 595 600 605

Ala Arg Arg Leu Ser Gly Pro Leu His Thr Leu Gly Ile Pro Pro Gly 610 620

Pro Asn Cys Thr Pro Ala Gln Gly Ser Arg Trp Pro Met Glu Lys Lys 625 630 635 640

Arg Arg Pro Ser Ala Leu Glu Ala Asp Ser Pro Met Ala Ser Lys
645 650 655

Arg Gly Thr Lys Arg Gln Arg Gln Ser Phe Leu Pro Cys Leu Arg Arg 660 665 670

Gly Ser Leu Pro Asp Thr Gln Pro Ser Gln Gly Pro Ser Thr Pro Lys 675 680 685

Gly Glu Arg Ala Ser Ser Pro Cys His Ser Pro Arg Val Cys Pro Ala 690 695 700

Thr Val Ile Lys Ser Arg Val Pro Leu Gly Pro Ser Ala Met Gln Asn 705 710 715 720

Cys Ser Thr Pro Leu Ala Leu Pro Thr Arg Asp Leu Asn Ala Thr Phe
725 730 735

Asp Leu Ser Glu Glu Pro Pro Ser Lys Pro Ser Phe His Glu Cys Ile 740 745 750

Gly Trp Asp Lys Ile Pro Gln Glu Leu Ser Arg Leu Asp Gln Pro Phe 755 760 765

Ile Pro Arg Ala Pro Val Pro Leu Phe Thr Met Lys Gly Pro Lys Pro 770 775 780

Thr Ser Ser Leu Pro Gly Thr Ser Ala Cys Lys Lys Lys Arg Val Ala
785 790 795 800

Ser Ser Ser Val Ser His Gly Arg Ser Arg Ile Ala Arg Leu Pro Ser 805 810 815

Ser Thr Leu Lys Arg Pro Ala Gly Pro Leu Val Leu Pro Glu Leu Pro 820 825 830

Leu Ser Pro Leu Cys Pro Ser Asn Arg Arg Asn Gly Lys Asp Leu Ile 835 840 845

Arg Val Gly Arg Ala Leu Ser Ala Gly Asn Gly Val Thr Lys Val Ser 850 855 860

<210> 3

<211> 1014

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: HsKip3a fragment

<220>

<223> Nucleotide sequence encoding motor domain fragment of HsKip3a (Figure 2).

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cccgatggag ggttccctgg cctgaaatgg ggtggcaccc atgatggccc caagaagaag 180
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gtgtttgcct acggggccac cggggctggg aagacacaca ccatgctggg aagggagggg 360
gaccccggca tcatgtacct gaccaccgtg gaactgtaca ggcgcctgga ggcccgccag 420
caggagaagc acttcgaggt gctcatcagc taccaggagg tgtataatga acagatccat 480
gacctcctgg agcccaaggg gccccttgcc atccgcgagg accccgacaa gggggtggtg 540
gtgcaaggac tttctttcca ccagccagcc tcagccgagc agctgctgga gatactgacc 600
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catgccatct tccagatctt tgtgaagcag caggaccggg ttccaggact gacccaggct 720
gtccaggtgg ccaagatgag cctgattgac ctggctggct cagagcgggc atccagcacc 780
catgcgaagg gggagcggct gcgggagggg gccaacatca accgctctct gctggcgctc 840
atcaacgtcc tcaatgcctt ggccgatgca aagggccgca agacccatgt gccctaccgg 900
gacagcaaac tgacccgcct gctcaaagac tccctcgggg gcaactgccg cacagtgatg 960
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<210> 4
<211> 338
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: HsKip3a fragment.
<220>
<223> Amino acid sequence of the motor domain fragment
      of HsKip3a (Figure 3).
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                                     10
Glu Leu Asp Ser Gln Arg Arg Pro Val Val Gln Val Val Asp Glu Arg
                                 25
Val Leu Val Phe Asn Pro Glu Glu Pro Asp Gly Gly Phe Pro Gly Leu
Lys Trp Gly Gly Thr His Asp Gly Pro Lys Lys Gly Lys Asp Leu
Thr Phe Val Phe Asp Arg Val Phe Gly Glu Ala Ala Thr Gln Gln Asp
Val Phe Gln His Thr Thr His Ser Val Leu Asp Ser Phe Leu Gln Gly
Tyr Asn Cys Ser Val Phe Ala Tyr Gly Ala Thr Gly Ala Gly Lys Thr
His Thr Met Leu Gly Arg Glu Gly Asp Pro Gly Ile Met Tyr Leu Thr
                                                 125
Thr Val Glu Leu Tyr Arg Arg Leu Glu Ala Arg Gln Gln Glu Lys His
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Phe Glu Val Leu Ile Ser Tyr Gln Glu Val Tyr Asn Glu Gln Ile His

150

Asp Leu Leu Glu Pro Lys Gly Pro Leu Ala Ile Arg Glu Asp Pro Asp
165 170 175

Lys Gly Val Val Gln Gly Leu Ser Phe His Gln Pro Ala Ser Ala 180 185 190

Glu Gln Leu Glu Ile Leu Thr Arg Gly Asn Arg Asn Arg Thr Gln
195 200 205

His Pro Thr Asp Ala Asn Ala Thr Ser Ser Arg Ser His Ala Ile Phe 210 215 220

Gln Ile Phe Val Lys. Gln Gln Asp Arg Val Pro Gly Leu Thr Gln Ala 225 230 235 240

Val Gln Val Ala Lys Met Ser Leu Ile Asp Leu Ala Gly Ser Glu Arg
245 250 255

Ala Ser Ser Thr His Ala Lys Gly Glu Arg Leu Arg Glu Gly Ala Asn 260 265 270

Ile Asn Arg Ser Leu Leu Ala Leu Ile Asn Val Leu Asn Ala Leu Ala 275 280 285

Asp Ala Lys Gly Arg Lys Thr His Val Pro Tyr Arg Asp Ser Lys Leu 290 295 300

Thr Arg Leu Leu Lys Asp Ser Leu Gly Gly Asn Cys Arg Thr Val Met 305 310 315 320

Ile Ala Ala Ile Ser Pro Ser Ser Leu Thr Tyr Glu Asp Thr Tyr Asn
325 330 335

Thr Leu

<210> 5

<211> 383

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: HsKip3a fragment

<220>

<223> Amino acid sequence of HsKip3a fragment used in the ATPase assay (Figure 4).

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Met Ala Val Glu Asp Ser Thr Leu Gln Val Val Val Arg Val Arg Pro 1 5 10 15

Pro Thr Pro Arg Glu Leu Asp Ser Gln Arg Arg Pro Val Val Gln Val
20 25 30

Val Asp Glu Arg Val Leu Val Phe Asn Pro Glu Glu Pro Asp Gly Gly

40

Phe	Pro 50	Gly	Leu	Lys	Trp	Gly 55	Gly	Thr	His	Asp	Gly 60	Pro	Lys	Lys	Lys
Gly 65	Lys	Asp	Leu	Thr	Phe 70	Val	Phe	Asp	Arg	Val 75	Phe	Gly	Glu	Ala	Ala 80
Thr	Gln	Gln	Asp	Val 85	Phe	Gln	His	Thr	Thr 90	His	Ser	Val	Leu	Asp 95	Ser
Phe	Leu	Gln	Gly 100	Tyr	Asn	Сув	Ser	Val 105	Phe	Ala	Tyr	Gly	Ala 110	Thr	Gly
Ala	Gly	Lys 115	Thr	His [.]	Thr	Met	Leu 120	Gly	Arg	Glu	Gly	Asp 125	Pro	Gly	Ile
Met	Tyr 130	Leu	Thr	Thr	Val	Glu 135	Leu	Tyr	Arg	Arg	Leu 140	Glu	Ala	Arg	Gln
Gln 145	Glu	Lys	His	Phe	Glu 150	Val	Leu	Ile	Ser	Tyr 155	Gln	Glu	Val	Tyr	Asn 160
				165			Glu		170					175	
Glu	Asp	Pro	Asp 180	Lys	Gly	Val	Val	Val 185	Gln	Gly	Leu	Ser	Phe 190	His	Gln
Pro	Ala	ser 195	Ala	Glu	Gln	Leu	Leu 200	Glu	Ile	Leu	Thr	Arg 205	Gly	Asn	Arg
Asn	Arg 210	Thr	Gln	His	Pro	Thr 215	Asp	Ala	Asn	Ala	Thr 220	Ser	Ser	Arg	Ser
His 225	Ala	Ile	Phe	Gln	Ile 230	Phe	Val	Lys	Gln	Gln 235	Asp	Arg	Val	Pro	Gly 240
Leu	Thr	Gln	Ala	Val 245	Gln	Val	Ala	Lys	Met 250	Ser	Leu	Ile	Asp	Leu 255	Ala
Gly	Ser	Glu	Arg 260	Ala	Ser	Ser	Thr	His 265	Ala	Lys	Gly	Glu	Arg 270	Leu	Arg
Glu	Gly	Ala 275	Asn	Ile	Asn	Arg	Ser 280	Leu	Leu	Ala	Leu	Ile 285	Asn	Val	Leu
Asn	Ala 290	Leu	Ala	Asp	Ala	Lys 295	Gly	Arg	Lys	Thr	His 300	Val	Pro	Tyr	Arg
Asp	Ser	Lys	Leu	Thr	Arg 310	Leu	Leu	Lys	Asp	Ser 315	Leu	Gly	Gly	Asn	Cys 320
Arg	Thr	Val	Met	Ile 325	Ala	Ala	Ile	Ser	Pro 330	Ser	Ser	Leu	Thr	Tyr 335	Glu
Asp	Thr	Tyr	Asn 340	Thr	Leu	Lys	Tyr	Ala 345	Asp	Arg	Ala	ГÀа	Glu 350	Ile	Arg
Leu	Lys	Gly	Asn	Ser	Lys	Leu	Glu	Gly	Lys	Pro	Ile	Pro	Asn	Pro	Leu

Leu Gly Leu Asp Ser Thr Arg Thr Gly His His His His His His 370 375 380

<210> 6
<211> 1152
<212> DNA
<213> Artificial Sequence

(213) Attititud bequene

<220>

<223> Description of Artificial Sequence: HsKip3a
 fragment

<220>

<223> Nucleotide sequence of HsKip3a fragment used in ATPase assay (Figure 5).

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